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I (ACC. NM		CAGTGCTCAG CATTACGŁAA GAATGGCTTT CAGTGCTCGG CCTTGCGCAA AAACGGCTTC	ATGLAAGATC GTCGAGATGT CTACTTCGAA ATGCAAAATA GTGGAGATGT CAACTTCCAA	CCAAGGTCCA TCTGGTTGGT ATTGACATCT CCAAGGTTCA CCTTGTTGGA ATTGATATTT	GAAATATGAA GATATCTGCC CGTCAACTCA ŁAATATGGAT AAAATATGAA GATATTTGTC CTTCTACTCA CAACATGGAT	TCAAAAGGAA TGACTtcCAG CTGATTGGCA TCCAGGATGG TTAAGAGAAA TGATTatCAA CTGATATGCA TTCAAGATGG	ACAGCGGGGA AAACTGGTGA		GCTGT.cTGCC / CATGT.gTGCA /	AATAa ATAA-
Human erro	ATGGCAGATG ACTTGGACTT ATGGCAGACG AAATTGATTT	CTTCCCAATG TTACCCTATG	AAGGCCGGCC AAGGACGACC	CAcGGCCACG CAtGGTCATG	GAAATATGAA AAAATATGAA	TCAAAAGGAA TTAAGAGAAA	CTGCTccagG ACAGCGGGGA CTGCTgacaG AAACTGGTGA	AGACCTTGGC AAGGAGATTG TGAACTAGGC AAAGAAATAG	TGATCACGGT AGGTGTCTGT	GCCatgGCAA AATAa CCCtGCAA ATAA-
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## FIG.59A

(0320)

SKTGK SKTGK	DGYLS DGYLS	AVAIK AVAIK	
VVLKGwPCKI VEMS/ VVLKGrPCKI VEMST	VPNIKRNDFQ LIGIC VPNIKRNDYQ LICIC	EEILITLLSA MTEEA EDVQVSVMCA MSEEY	
TFPM QCSALRKNGF TYPM QCSALRKNGF	KKYE DICPSTHNMD V KKYE DICPSTHNMD V	GDLG KEIEQKYDCG I GELG KEIEGKYNAG I	
MADDLDFETG DAGASATFPM QCSALRKNGF VVLKGWPCKI VEMSASKTGK I MADEIDFTTG DAGASSTYPM QCSALRKNGF VVLKGrPCKI VEMSTSKTGK	HGHAKVHLVG IDIFTGKKYE DICPSTHNMD VPNIKRNDFQ LIGIQDGYLS HGHAKVHLVG IDIFTGKKYE DICPSTHNMD VPNIKRNDYQ LICIQDGYLS	LLQDSGEVPE DLRLPEGDLG KEIEQKYDCG EEILITLLSA MTEEAAVAIK LLTETGEVRE DLKLPEGELG KEIEGKYNAG EDVQVSVMCA MSEEYAVAIK	amak pck-
	51 F	101 101 L	151 a 151 p
eIF5a1 eIF5a2	eIf5a1 eIF5a2	eIf5a1 eIF5a2	eIf5al eIF5a2

FIG.59B